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Aquatic tri-trophic standardized microcosm TriCosm

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A standardized tri-trophic small-scale system (TriCosm) for the assessment of stressor induced effects on aquatic community dynamics

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ABSTRACT

Chemical impacts on the environment are routinely assessed in single-species tests. They are employed to measure direct effects on non-target organisms but indirect effects on ecological interactions can only be detected in multi-species tests. Micro- and mesocosms are more complex and environmentally realistic, yet, they are less frequently used for environmental risk assessment because resource demand is high while repeatability and statistical power are often low. Test systems fulfilling regulatory needs (i.e. standardization, repeatability and replication) and the assessment of impacts on species interactions and indirect effects are lacking. Here we describe the development of the TriCosm, a repeatable aquatic multi-species test with three trophic levels and increased statistical power. High repeatability of community dynamics of three interacting aquatic populations (algae, *Ceriodaphnia*, *Hydra*) was found with an average coefficient of variation of 19.5% and the

ability to determine small effect sizes. The TriCosm combines benefits of both single-species tests (fulfillment of regulatory requirements) and complex multi-species tests (ecological relevance) and can be used, for instance at an intermediate tier in environmental risk assessment. Furthermore, comparatively quickly generated population and community toxicity data can be useful for the development and testing of mechanistic effect models.

KEYWORDS

ecological risk assessment, aquatic invertebrates, population-level effects, food chain, aquatic microcosm, multi-species testing

INTRODUCTION

The thorough assessment of environmental risks is essential for chemicals that could potentially be released into the environment. Agricultural pesticides, for instance, are used to enhance crop production but due to their toxic nature they may have negative effects on organisms other than the targeted species (Benton et al. 2007; Rockström et al. 2009; Beketov et al. 2013; Stehle and Schulz 2015).

The risks of pesticides to non-target organisms are routinely assessed in i) simple single-species tests at lower tiers and, if lower tier assessments raise concern, in ii) complex microcosms or mesocosms (European Food and Safety Authority (EFSA) 2013). However, systems that bridge the gap between the two alternatives to an intermediate level of complexity are lacking. The former require less effort and rapidly deliver large amounts of highly repeatable data on the performance of individual non-target organisms. The information obtained is, however, often of low ecological relevance as it is not directly relevant at the population and community level (Fleeger et al. 2003; Liebig et al. 2008). In

contrast, micro-/mesocosms are environmentally more realistic, yet, they are less frequently used in environmental risk assessment. Unlike single-species tests, they are resource, time and effort demanding. A variety of ecologically interacting factors can rapidly lead to divergent system dynamics and increase the variance between replicates impeding the understanding of dose-response relationships (Landis et al. 1997).

The importance of integrating environmental complexity into testing approaches has been acknowledged as a priority for the assessment of chemical safety (Landis et al. 1997; Landis 2003; Bednarska et al. 2013; Scientific Committee on Health and Environmental Risks (SCHER) et al. 2013). Chemical exposure could trigger indirect effects through interactions with the environmental context such as the hydrological regime (Stampfli et al. 2013), temperature (Moe et al. 2013), food quality (Campos et al. 2014) or other organisms (Del Arco et al. 2015; Viaene et al. 2015); indirect effects have important implications for the sensitivity of communities (Fleege et al. 2003).

Organisms living in a contaminated environment may be pushed towards the boundaries of their ecological niche and become more susceptible to additional stressors (Van Straalen 2003; Bednarska et al. 2013). Food chain processes, such as competition for food and altered predation were shown to be particularly relevant to determine the magnitude of toxic effects (Heugens et al. 2001; Bednarska et al. 2013; Kattwinkel et al. 2015). For instance, the no observed effect concentration (NOEC) of the herbicide prometryn to ciliates was found to be approximately 145 times lower in a bi-trophic microcosm compared to single-species tests. The lower threshold was likely caused in response to an indirect and toxicant induced reduction of food (Liebig et al. 2008). Intraspecific competition can also change the sensitivity to pesticides (Foit, Kaske, and Liess 2012; Viaene et al. 2015) and indirectly altered predation rates can lead to cascading effects on other trophic interactions and ecosystem functions (Englert et al. 2012; Agatz et al. 2014; Viaene et al. 2015). Multi-species

testing using environmentally more relevant approaches, i.e. at the population and community level is clearly needed to assess indirect toxicant effects such as shifts in ecological interactions (Fleege et al. 2003; Benton et al. 2007).

The necessity towards an inclusion of ecological interactions in chemical impact testing was described 10 years ago when a review on 14 years of pesticide studies in freshwater test systems was published (Relyea and Hoverman 2006). At the time, the authors found only 133 studies with at least two potentially interacting species of which only 17 studies focused on three trophic levels with producers, herbivores and carnivores.

Yet, microcosms that describe impacts on populations and/or communities in systems smaller than 10 L are rare (Metcalf et al. 1971; Daam and Van Den Brink 2007; Liebig et al. 2008; Englert et al. 2012; Foit, Kaske, and Liess 2012; Dolciotti et al. 2014; Del Arco et al. 2015; Viaene et al. 2015). Mostly they were used to focus on impacts on intra- or interspecific competition (one trophic level) (Foit, Kaske, and Liess 2012; Dolciotti et al. 2014; Del Arco et al. 2015; Viaene et al. 2015) or on consumer-resource relationships (two trophic levels) with herbivore-producer (Daam and Van Den Brink 2007) or predator-prey interactions (Barry and Davies 2004; Liebig et al. 2008; Englert et al. 2012). Tri-trophic systems are frequently used in terrestrial research, for example in plant-herbivore-parasite systems (Bredeson et al. 2015; Uhl et al. 2015) but few small test systems exist to assess direct and indirect impacts at the population and community level in the aquatic environment. Test formats include simulations of microbial detritus food chains (producer-consumer-decomposer (Fuma et al. 2000; Dawoud et al. 2017)) and producer-consumer communities with either invertebrate predator (Barry and Davies 2004) or vertebrate predator (Metcalf et al. 1971). Microbial tests were often conducted in culture flasks (250 ml) (Fuma et al. 2000; Dawoud et al. 2017) and small macroinvertebrate community tests were performed in

systems of few litres, for example in 10 L (Barry and Davies 2004) and 7 L systems (Metcalf et al. 1971).

Still, single-species systems appear convenient because they fulfill the regulatory needs for international standardization of test procedures, comparability of effect data, repeatability and replication (Liebig et al. 2008). Standardized and repeatable multispecies systems of intermediate complexity that bridge the simplicity of single species tests and the complexity of microcosms, yet fulfill regulatory requirements, are rare. To our knowledge, there is only one standardized microcosm (Taub 1989) available that falls into this category. The aquatic system was registered for pesticide testing (American Society for Testing of Materials (ASTM) 2011) and effects on two trophic levels covering ten primary producer and five primary consumer species can be assessed. It is, however, rarely used for standardized effect assessment, perhaps due to its relative complexity and the lack of mechanistic understanding of the interactions between species involved.

We developed a new test system with species interacting across three trophic levels and increased statistical power (i.e. standardization and low replicate variability). The system was designed to be cost-effective, rapid, repeatable with well understood population dynamics to i) allow the detection of small changes in population dynamics due to direct and indirect interactions, and ii) link observed effects to known system processes. Here we describe the standardized aquatic tri-trophic microcosm (hereafter TriCosm) focusing on system design and variability in the control treatment.

MATERIALS AND METHODS

Test organisms

The TriCosm comprises populations of the green alga *Pseudokirchneriella subcapitata*, the cladoceran *Ceriodaphnia dubia* and the cnidarian *Hydra viridissima*. This

dynamic food-chain is subject to fluctuating but predictable changes in food supply and intraspecific competition and is interconnected through consumer-resource relationships. The species were chosen based on their rapid life cycles and their sensitivity to toxicants. The green alga *P. subcapitata* and the cladoceran *C. dubia* are routinely used for tests in the regulatory risk assessment framework (Organization for Economic Co-operation and Development (OECD) 2004; OECD 2006; OECD 2012).

P. subcapitata stock cultures were obtained from the Culture Collection of Algae and Protozoa (CCAP, Scotland, UK) and used to initiate a culture line prior to each study and cultured in OECD media (OECD 2006). *C. dubia* were obtained from Unilever (Safety and Environmental Assurance Centre, Bedford, UK) and cultured as age specific cultures in moderately hard, synthetic freshwater (United States Environmental Protection Agency 2002). They were fed five times per week with a suspension of yeast, cerophyl[®] and trout-chow (~3.5 ml) and *P. subcapitata* (~11 x 10⁷ cells/day) (United States Environmental Protection Agency 2002). *H. viridissima* were obtained from the Department of Evolutionary Zoology (University of Debrecen, Hungary), cultured in modified T82MV medium (modified after ASTM E1366-11 2011, Table SI 1, Table SI 2) and fed with newly hatched *Artemia salina* three times per week ad libitum. Both animal cultures were kept at 25 ± 1 °C and 12/12h light/dark cycle.

The TriCosm

TriCosms consist of Pyrex[®] crystallizing dishes (Sigma-Aldrich, UK) filled with 500 ml of T82MV medium (ASTM E1366-11 2011, Table SI 1, Table SI 2) that was determined as suitable for each species. The systems were covered with transparent watch glasses (diameter 125 mm; Sigma-Aldrich, UK) and positioned on an orbital laboratory shaker (Adolf Kuehner AG Switzerland, Type LS-W) set at 65 rpm throughout the test. The

experiments were set up for 21 days at 25 ± 1 °C, 12/12h light/dark, 1100 lux at the water surface with cool white fluorescent light tubes 58.5 W (approx. 1.3 m above the test vessels). TriCosms were started with *P. subcapitata* cells from a culture in logarithmic growth phase, *C. dubia* neonates (< 24h age) from the third or fourth brood of cultured mothers and *H. viridissima* without visible buds (≤ 2 d age). Water parameters (pH, dissolved O₂) and animal and algal populations were monitored throughout the test duration two, three and five times per week, respectively.

Monitoring of the populations

The systems were placed on an orbital shaker and slow shaking kept the algal cells suspended. Only suspended algae were measured and no stirring was necessary before sampling as preliminary studies showed significant correlation between suspended and total algal concentrations (cells/ml) ($r_s = 0.98$, $p < 0.01$, $n = 90$, **Figure SI 1**). In-vivo fluorescence activity of water subsamples (5 x 200 µl) was measured with a plate reader (Tecan® Infinite 200 PRO, settings **Table SI 3**) to determine the algal concentration (cells/ml).

C. dubia and *H. viridissima* were monitored with non-invasive methods to avoid impacts on population dynamics and counted by eye three times per week. *C. dubia* were visually grouped in two age-classes, juveniles and adults based on their dimensional similarity with individuals in cultures aged younger or older than 4 d. All manual counts were repeated until count differences did not exceed 20% of the lower value.

Assessing a suitable community composition

Tests with different setups were performed to optimize replicate variability, test duration, addition times and densities for each species. A full factorial design for density and timing was not feasible due to a too high number of possible combinations. Hence,

preliminary tests were performed to determine which algae-grazer combination in terms of organism abundance would prevent both algal blooms and the death of grazers due to starvation. No preliminary tests were done to determine the impact of *Hydra* predation on *C. dubia* numbers prior to the test outlined in Table 1. Two organism densities and different addition times were chosen based on preliminary testing and four different setups were conducted simultaneously (Table 1). *C. dubia* were added on the same day as the green algae in all experiments, except for setup 1 where grazers were added 1 day later to allow short acclimation of the algae to test conditions. Dependent on food concentrations, *C. dubia* matured later in setups 1, 2 than in setups 3, 4 hence *H. viridissima* were introduced to the systems 5 and 4 days, respectively, after *C. dubia* were added. The predators were added only once the grazers started reproducing to prevent variable numbers of *C. dubia* reproducers and neonates early on in the systems that could lead to noticeable impacts on community dynamics and replicate variability. Replicate numbers differed between setups 1, 2 and 3, 4 due to space constraints on the shaker platform.

Validation of an optimal experimental setup

The coefficient of variation (CV) was calculated as a standardized measure of variance between replicates. It was expressed as a percentage and indicates the magnitude of the standard deviation in comparison to the mean. Thus, if the inherent variability between replicates is large compared to the size of the measured endpoint (e.g. animal abundance) a significant treatment effect could only be detected if the response was very large (Sanderson 2002).

We computed the CVs for the algal concentration (cells/ml) and the total number of *C. dubia* and *H. viridissima* on every sampling day. The values were then compared within and between setups to monitor replicate variation over time and to determine an optimal setup

in terms of low variance. The experimental setup with the lowest replicate variability (Experiment 1) was determined and repeated (Experiment 2) to assess the reproducibility of system dynamics and replicate variability.

Population dynamics and interactions

In the interacting system, the intermediate trophic layer is directly affected by both variations of food availability and predation strength, while indirect effects between the bottom and top trophic level regulate a bottom up or top down controlled system. A comparison of algal and grazer dynamics between systems where grazers are subject to i) variations of food but not to predation and ii) a combination of food limitation and predation can thus yield information on species interaction strength and whether the system is controlled by bottom up or top-down effects. Hence, we performed additional tests with i) only algae (Experiment 2A, n = 8) and ii) algae and grazers (Experiment 2B, n = 8). The experiments were carried out simultaneously to Experiment 2 and according to the experimental conditions and test setup used as for Experiments 1 and 2 (see Table 1 for details). Experiments 2A and 2B were then compared to determine direct impacts of grazers on algal dynamics and Experiments 2 and 2B were compared to assess i) indirect effects of predators on algal growth and ii) direct effects of predators on *C. dubia* population trajectories.

Statistical analyses

An a priori power analysis was performed to estimate minimum detectable response sizes between control and treated TriCosm populations and increase the reliability and transparency of the derived endpoints (EFSA 2013). The minimum detectable difference (MDD), i.e. the size of a variation between sample averages required to be detected as

significantly different, is dependent on the chosen Type I error value α , the number of replicates employed and on the inherent variance such as replicate variability and/or sampling error (Brock et al. 2015). Here, we computed the MDD using the CVs assuming similar variance among controls and treatments. We hypothesized the use of 8 replicates and estimated the sensitivity of the TriCosm to reveal chemical effects for each population and at each sampling point. The MDD was calculated as described by Brain et al. (2005) from Sokal and Rohlf (1995):

$$\text{MDD} = \frac{\sqrt{2} (t_{\alpha,v} + t_{\beta,v}) \times \text{CV}}{\sqrt{n}}$$

where $t_{\alpha,v}$ and $t_{\beta,v}$ are the t-values for α and β set to 0.05 and 0.2, respectively, for a confidence level of 95% and a power of 80% at v degrees of freedom. CV is the coefficient of variation and n is the number of replicates used. The degrees of freedom were computed as $v = k(n-1)$ and the number of groups k was set to 2, e.g. to compare each treatment to the control. The calculated MDDs were compared to MDD classes as proposed by the European Food Safety Authority (EFSA) that grouped MDD sizes into five classes and described the likely ability of effect detection (Class 0: MDD > 100% = no effect detection, Class I: MDD 90 – 100% = only large effects, Class II: MDD 70 – 90% = large to medium effects, Class III: MDDs 50 – 70% = medium effects, Class IV: MDD < 50% small effects) (EFSA 2013).

To assess species interactions between system components, population dynamics were compared graphically between experiments and significant differences were assumed where 95% confidence intervals did not overlap.

RESULTS

Optimizing the experimental setup

The community dynamics (Figure 1) and the coefficients of variation differed among the four setups (Figure 2) and over time (Figure SI 2). In general, the algal concentration (cells/ml) peaks were followed by *C. dubia* abundance peaks and a constant increase of *H. viridissima* populations. The highest *C. dubia* peak 172 (± 10) individuals (mean \pm 95% confidence interval range) was found in Setup 1 on day 14 following an algal peak on day 6 when an average of $6.86 (\pm 0.64) \times 10^5$ cells/ml was measured (Figure 1A). The highest algal peak of $11.51 (\pm 5.59) \times 10^5$ cells/ml appeared in Setup 3 on day 11 with increasing variance in terms of organism numbers between replicates over time (Figure 1C). *H. viridissima* populations showed steady growth during the test duration and increased in numbers by an average of 42 (± 6), 45 (± 8), 72 (± 18) and 54 (± 15) individuals in Setup 1, 2, 3 and 4, respectively (Figure 1A - D). Final counts differed due to different addition numbers, timings and food availability (*C. dubia* abundances), however, *Hydra* populations showed the smallest replicate variance when compared to algae and *C. dubia* (Figure 2). The CVs of all test variables in each setup increased over time (Figure SI 2) with setup 3 showing the highest replicate variability (with the exception of *H. viridissima*) and setup 1 showing the lowest average CV (with the exception of *H. viridissima*) (Figure 2). High CVs observed in setup 2 - 4 indicated reduced ability to detect treatment related system alterations. Therefore we selected setup 1 (Experiment 1) as most appropriate setup procedure (Protocol SI 4) and repeated the test (Experiment 2) to evaluate the repeatability of the system.

Validation of the test setup

The population dynamics of experiment 1 and the repeated experiment 2 were similar (Figure 3). The algal populations peaked on day 6 and day 5 with average algal concentrations of $6.86 (\pm 0.64) \times 10^5$ and $8.83 (\pm 0.90) \times 10^5$ cells/ml in experiments 1 and 2, respectively (Figure 3A). The highest *C. dubia* abundance was measured 8 days after the algal peaks in both cases. Grazer numbers declined 1 day earlier in experiment 2 and individual counts were lower due to smaller juvenile numbers of $147 (\pm 10)$ and $97 (\pm 24)$ juveniles in experiment 1 and 2, respectively (Figure 3B). The dynamics of *H. viridissima* populations were similar between experiments (Figure 3C) but experiment 1 showed a slightly steeper population increase with a larger final population of $45 (\pm 6)$ and $33 (\pm 9)$ individuals in experiment 1 and 2, respectively. Due to slightly different sampling frequencies, we computed 15, 11 and 8 CV values in experiment 1 and 14, 9 and 7 CVs in experiment 2 for algal concentrations (cells/ml), *C. dubia* and *H. viridissima*, respectively. As observed for population dynamics, replicate variance was similar between populations in both experiments (Figure SI 3). The CVs of algal concentrations (cells/ml) increased by day 7, 14 and 21 to an average of 9, 26 and 26% in experiment 1 and 10, 25 and 47% in experiment 2, respectively. The CVs calculated for *C. dubia* populations increased from 9 to 15 and 33% in experiment 1 and from 4 to 23 and 24% in experiment 2. *H. viridissima* were added on day 6, so the replicate variability was 18, 18% and 17, 29% by day 14 and 21 in experiment 1 and 2, respectively.

Population dynamics and interactions

Significant reductions of algal concentrations (cells/ml) by 33.4% were found on the first sampling day after *C. dubia* addition, on day 4. An average algal concentration of $9.18 (\pm 0.48) \times 10^5$ cells/ml was found in Experiment 2A (only algae), while grazed algae in Experiment 2B (algae and grazers) reached an abundance of $6.11 (\pm 0.43) \times 10^5$ cells/ml by

day 4 and stayed significantly lower throughout the experimental duration (Figure 4A). Grazed algal concentrations in systems with and without predator (Experiments 2 and 2B, respectively) showed similar trajectories until day 14. After day 14, algal concentrations in Experiment 2B stayed moderately constant with an average of $2.12 (\pm 0.21) \times 10^5$ cells/ml until day 20. On the contrary, algal concentrations in Experiment 2 (grazers and predators) increased to $7.63 (\pm 0.37) \times 10^5$ cells/ml by day 20 exceeding average algal abundances of Experiment 2B by 75.0% (Figure 4B).

Grazer population dynamics were similar in Experiment 2 and 2B until day 6 when predators were added to Experiment 2 (Figure 4C). Population numbers peaked in both experiments on day 13 but steeper population growth curves in Experiment 2B lead to an on average 39.1% larger peaking population of $210 (\pm 21)$ individuals when compared to $128 (\pm 25)$ individuals in Experiment 2. A similarly steeper population decrease in experiment 2B resulted in similar total *C. dubia* counts of $36 (\pm 9)$ and $37 (\pm 10)$ individuals in systems without and with predators, respectively, by day 20. Population dynamics of total grazer numbers largely reflected the trajectories of *C. dubia* juveniles that rapidly increased until day 13 to $175 (\pm 23)$ and $107 (\pm 23)$ individuals constituting 88.3% and 90.1% of the total *C. dubia* population in experiments 2B and 2, respectively. By day 20, juvenile numbers dropped to $6 (\pm 4)$ and $15 (\pm 6)$ individuals while adult grazers showed a moderate but constant increase throughout the test and constituted 84.9% and 59.7% of the total *C. dubia* populations in experiments without and with predators, respectively (Figure 4D).

Statistical analyses

Minimum detectable differences (MDDs) of hypothetical TriCosm exposures were calculated according to control variance and were similar between experiments 1 and 2. The TriCosm became less sensitive over time as replicate variation and MDDs increased. When

variances between controls and treatments are similar, the TriCosm is estimated to be sufficiently sensitive to identify differences of 12% (± 4), 36% (± 7) and 50% (± 17) for *P. subcapitata* and 9% (± 7), 31% (± 4) and 38% (± 19) for *C. dubia* populations in week 1 (day 1 – 6), week 2 (day 7 – 13) and week 3 (day 14 – 21), respectively (averaged MDDs between Experiment 1 and 2). Averaged MDDs for *H. viridissima* were 25% (± 14) and 35% (± 7) in week 2 and 3, respectively (Figure 3). The MDDs for critical endpoints in the TriCosm can be assigned to MDD classes III (50 – 70%) and IV ($< 50\%$), indicating the ability to determine small and medium sized effects, respectively (EFSA 2013).

DISCUSSION

The assessment of chemical effects with single-species tests fulfills regulatory requirements; however, primary goals of protecting populations and ecosystems might not be adequately addressed. That is because information obtained at the individual level is often not ecologically relevant since there are neither directly proportionate relationships between direct and indirect effects nor amongst responses at the individual, population and community level. An understanding of impacts on interactions in ecologically relevant test settings is thus critical and a priority for chemical safety assessment as unexpected shifts in community profiles cannot be predicted in single-species tests (Fleege et al. 2003; Benton et al. 2007; Liebig et al. 2008; SCHER et al. 2013).

We designed the TriCosm as a rapidly cycling, tri-trophic system with a producer-herbivore-carnivore community of small size for the purpose of quick detection of chemical impacts on species interactions. Our system is comparatively smaller (0.5 L) than many other multi-trophic systems (Metcalf et al. 1971; Daam and Van Den Brink 2007; Foit, Kaske, Wahrendorf, et al. 2012; Dolciotti et al. 2014; Del Arco et al. 2015) and all system components exhibit rapid generation times so that treatment effects can be measured on

several generations and at different life stages during short test durations (21 days compared to 80 days (Metcalf et al. 1971) and 33 days (Barry and Davies 2004) in other tritrophic macroinvertebrate communities). Also the predator *Hydra* is a rapid reproducer with generation times of only three days under favourable conditions (Habetha et al. 2003). Chemical impacts on population dynamics can thus be detected not only at the producer-consumer level but also at a higher trophic level. The choice of a small and rapidly reproducing predator has further the advantage that it can be added at an early experimental stage (day 6) when compared to vertebrate predators that are often introduced shortly before test termination as they quickly consume remaining invertebrate preys (Metcalf et al. 1971; Harrass and Taub 1985).

All multi-species systems have ecologically interacting components that are not independent in statistical terms as they constantly adapt to changing conditions in a dynamic environment. In fact, it has been frequently reported that even though communities are set up identically as replicates, minor variations at the beginning and/or throughout the experiments (e.g. starting conditions or uneven sample removal) can quickly lead to the development of unique properties in each replicate (Landis et al. 1997; Sanderson 2002; Van Straalen 2003). Indeed, different population dynamics and replicate variability were observed in four different TriCosm setups and indicated strong sensitivity to starting conditions and interaction strength. The statistical quality (in terms of interpretability, reproducibility and replicability) of environmentally more realistic data obtained in multi-species tests is thus often reason for concern in the registration procedure of pesticides (Sanderson 2002).

The repeatability and reproducibility of the TriCosm were thus given major consideration during test development. Initial properties and sampling techniques were adjusted and confirmed as optimized when experiments conducted at different times showed low coefficients of variation (CVs) and high reproducibility of system dynamics.

Desynchronized population dynamics were observed between experiments that can be attributed to random fluctuations in test conditions (e.g. quality of the animals) and could occur even if procedures are standardized. For these reasons we assessed the repeatability by comparing CVs and not the total organism abundances. Nonetheless, a comparison of total abundances or derived variables (e.g. population growth rates) is also appropriate when chemical effects are assessed since differences between population trajectories are most likely and primarily due to chemical impacts rather than fluctuations of test conditions.

When the TriCosm is used for chemical effect assessment, two factors of major importance are i) the presence of interactions rather than the exact timing when these occur and ii) low CVs so that treatment responses can be interpreted with greater certainty and distinguished from unexplained sample variability (Sanderson 2002).

The ability to detect significant effects does depend on the magnitude of an effect but also on the ability of the test system to detect responses and that is in turn determined by the inherent variance among replicates. Test variables with coefficients of variation (CV) in the range of up to 30% have been theorized as acceptable and manageable in terms of practicality and costs (Kraufvelin 1998). According to a review (Sanderson 2002) that analyzed two decades of pesticide studies with micro/mesocosms, the values of CVs appear to be generally higher. The author reported an average of 45% (32% in smaller and less realistic indoor systems) with larger values in studies where animals were involved and an average use of 3.5 replicates. The average CV of 19.5% measured in the tri-trophic system on the contrary showed smaller variance and was determined with a higher number of replicates ($n = 8$).

The CVs were further used for the calculation of theoretically detectable minimum differences (MDDs) between controls and treatments under the assumption of similar variances. It is to be mentioned, however, that the variance could increase, decrease and/or remain similar in treated systems (Kraufvelin 1998; Sanderson 2002). A modification of the

number of replicates, groups or treatments, though, can decrease MDDs and allow the detection of desired effect sizes. Due to often large variability in micro-/mesocosms, EFSA may still regard endpoints with MDD classes I and II (70 – 100%) relevant but considers the exceeding of class II ideal (i.e. MDDs < 70%) (EFSA 2013). Most projected MDDs of critical endpoints in the aquatic system correspond to effect class IV (i.e. < 50%) (with exception of algae and grazers in week 3) and confirm the ability to reveal small toxicant induced effect sizes (EFSA 2013), distinguishing the TriCosm from other multi-trophic systems.

As expected, variations of population trajectories were observed as a result of interactions with other system components. Algal concentrations (cells/ml) and predation both directly impacted on the middle trophic layer while they indirectly impacted on the top and the bottom level, respectively. An initially small grazing pressure of juvenile *C. dubia* allowed algal populations to grow exponentially which in turn favored the development of grazer populations. As a consequence of an increasing grazing pressure by maturing and reproducing *C. dubia*, the algal concentrations dropped, yet the grazer population numbers further increased for approximately one week after food availability became limiting. The continuing population growth is attributable to a rise of juvenile numbers as adult *C. dubia* most likely matured eggs and stored energy before algal concentrations decreased. Peaking *C. dubia* populations thus coincided with lows of food availability and caused the decrease of grazer numbers. Algae stabilized and remained at relatively constant levels as concentrations were most likely too low to be further reduced if maximum grazer filtering rates were reached. Grazer population numbers decreased due to food shortage and independently of whether predators were present or not. While predation did not cause *C. dubia* populations to crash, it directly reduced grazer numbers, intraspecific competition among them and indirectly favored algal populations to recover. An increase of algal concentrations after

grazing release was, however, not observed in Experiment 1 where grazer populations reached larger abundances but decreased later and might be due to a different quality of neonates used to start the experiments. Algal populations in Experiment 1 were thus subject to a higher and prolonged grazing pressure impeding the recovery of algal abundances within the experimental duration. An indirect effect after grazing release by *Hydra* could, however, likely be expected if the test duration was prolonged. Bottom up and top down processes are thus both likely regulating population dynamics in the TriCosm. When the system is exposed to chemicals it will thus depend on the mode of action of the toxicant impacting on one or more trophic levels leading to direct, indirect or both effects on the trajectories of interacting populations.

The focus during system development was not on achieving a steady state community and impacts on resilience cannot be assessed, neither can system shifts be detected. Nonetheless, it can indicate the recovery potential of species after stressor removal and detect small changes in interactions as the system moves through a single cycle of the middle trophic layer. Ecological impacts of toxicants rapidly propagate in an interacting system and the grazer level is directly influenced by variations in food availability and predation. Toxicant impacts on the population dynamics of this critical and key trophic layer will therefore yield important information on the ecological relevance and protectiveness of data obtained in single-species tests. Population responses to combined stressor exposures, e.g. to toxicants, predation and/or food fluctuations, could be used to facilitate both the development and the testing of mechanistic effect models. Measured community responses in terms of individual abundance changes and population trajectories could be employed for the calibration and parameter fitting of ecological models. In turn, chemical effects on interactions within a simple freshwater community can be measured and quantified in the

TriCosm and provide empirical benchmarking to estimate and test model prediction accuracy and power.

There is no doubt that the complexity of the TriCosm community is low when compared to natural systems. But besides offering higher statistical power when compared to larger and / or more complex microcosms, the impacts on system processes can be quantified as interactions change. This makes it possible to assess the effects of environmental contaminants on i) species interactions, ii) indirect effects and iii) at the population and community level. An understanding of which and to what extent processes are affected may also give insights into responses of more complex systems (Benton et al. 2007; Daam and Van Den Brink 2007; Boonstra et al. 2011) .

CONCLUSION

The TriCosm is a novel aquatic test system and could be a tool to address shifts in ecological interactions. It suggests that a cost-effective approach of chemical environmental safety testing with more ecological relevance whilst being statistically powerful is feasible. It can provide important insights into chemical safety in multi-trophic systems and facilitate the development and testing of mechanistic effect models for environmental risk assessment. Even so, a careful examination of the replicability of the TriCosm both within and between laboratories with and without chemical exposure is needed.

Supplemental Data — The Supplemental Data are available on the Wiley Online Library at DOI: 10.1002/etc.xxxx

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Data availability — Data and calculation tools are available from the corresponding author (Verena.riedl@york.ac.uk).

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TABLES AND FIGURES

Table 1. The TriCosm community composition at the beginning of four test setups

	<i>P. subcapitata</i>		<i>C. dubia</i>		<i>H. viridissima</i>		Replicates
	Cells/ml	Day	Individuals	Day	Individuals	Day	
Setup 1	2 x 10 ⁴	0	10	1	3	6	8
Setup 2	2 x 10 ⁴	0	10	0	6	5	8
Setup 3	4 x 10 ⁴	0	10	0	6	4	7
Setup 4	4 x 10 ⁴	0	20	0	6	4	7

Figure 1 Algal concentrations (cells/ml) and total number of *C. dubia* and *H. viridissima* over 21 days. Shown are means ± 95 % confidence intervals in four test setups (A – D) (see Table 1 for details).

Figure 2 Coefficients of variation (%) of algal concentrations (cells/ml), total abundance of *C. dubia* and *H. viridissima* at each sampling event. Black horizontal lines indicate 95 % confidence intervals in setups 1 - 4.

Figure 3 Abundance of (A) *P. subcapitata*, (B) *C. dubia* and (C) *H. viridissima* at each sampling point over 21 days. Shown are means \pm 95% confidence intervals and minimum detectable differences (% MDD) below the x- axis of experiment 1 and experiment 2 (Exp 1, Exp 2).

Figure 4 Algal population trajectories compared between (A) ungrazed (green circles) and grazed (without predation, blue diamonds) systems and (B) grazed systems with (red stars) and without predation (blue diamonds). Population dynamics of *C.dubia* with (red stars) and without (blue diamonds) predation as (C) total individual number and (D) juveniles (continuous line) and adults (dotted line).

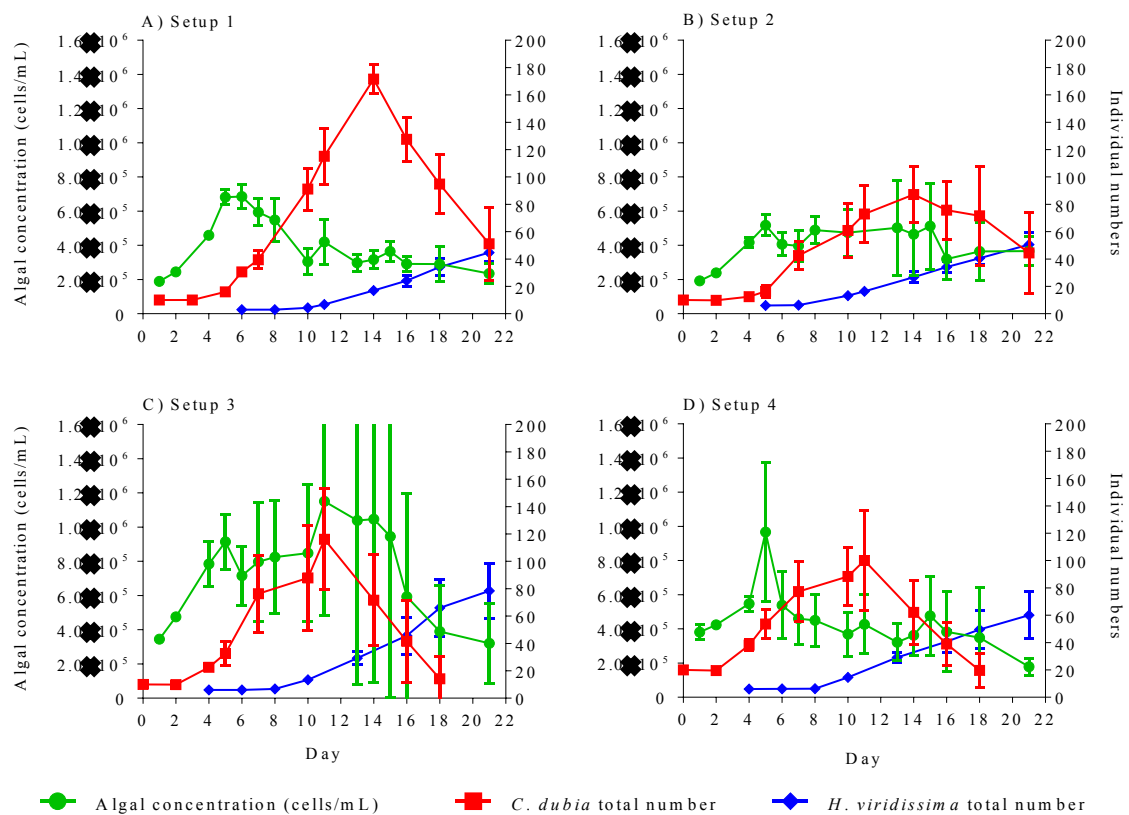


Figure 5

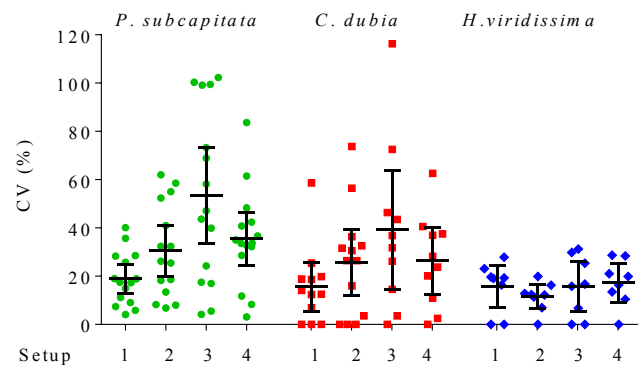


Figure 2

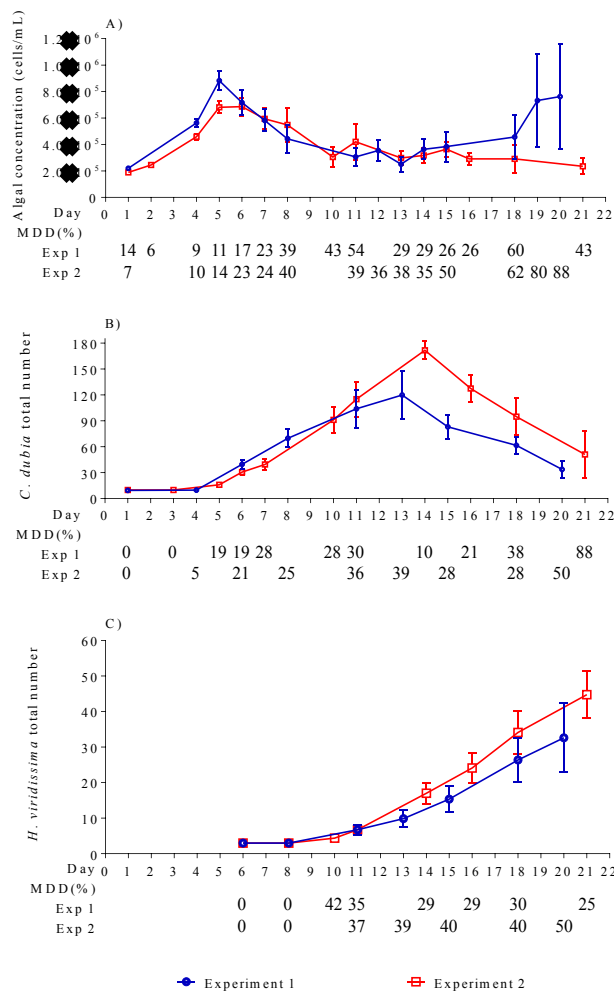


Figure 3

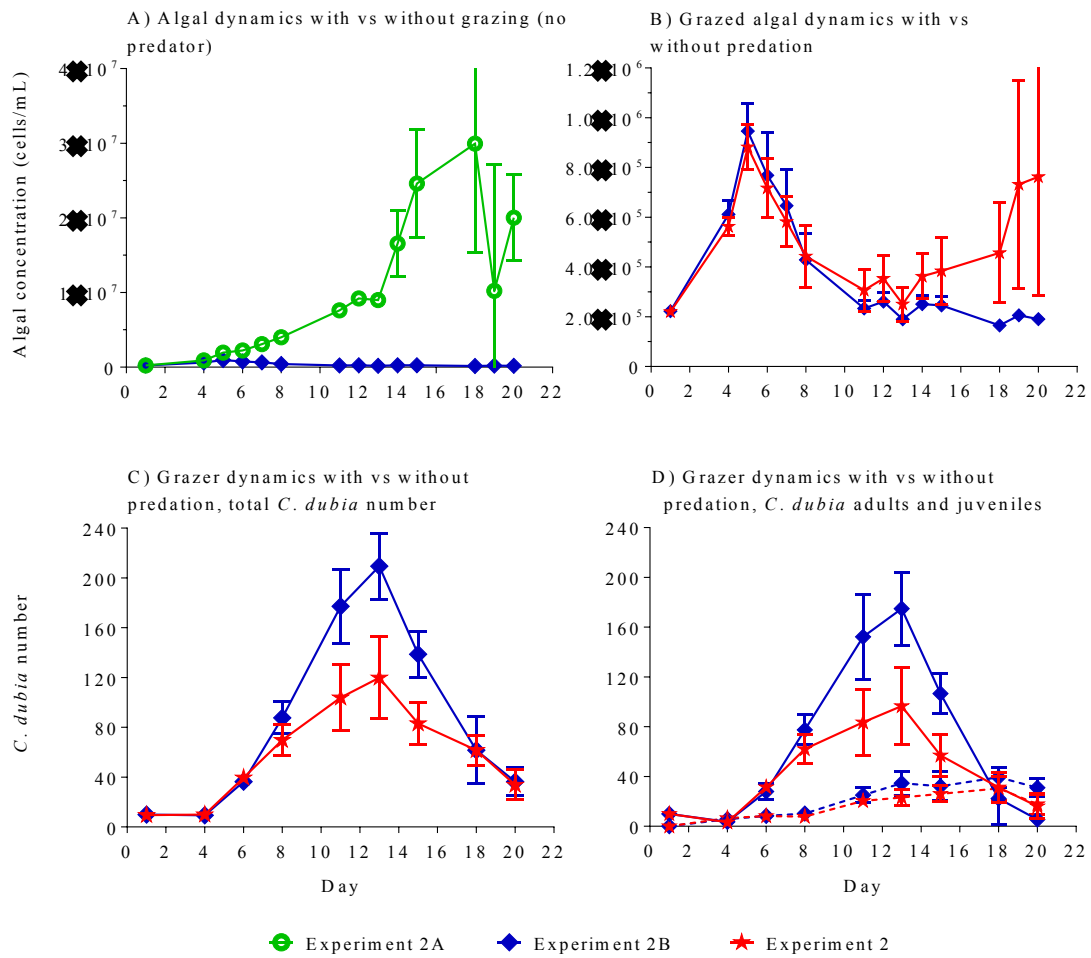


Figure 4